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121 LAGCTTGAMAVTCAOPTDVVKVRFOASIHLGPSRSDRKYSGTHDAYRTIARRECVRCLWK 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                               241 WERRIMSPROGESPLOCHIKMVAORGPTAFYKGFTPSFLELGSWINWHFTFEOLKRA JOD
241 WERRIMSPROGESPLOCHIMINION IN INTERCEPTATION OF THE JOD
241 WERRIMSPROGESPLOCHIMINAOSPTAKKEFPSFLAGSWINWHFYTFOLKRA JOD
                                                                                                                  Overy Match 100.0%; Score 1620; DB 1; Langth 312; Best Local Similarity 100.0%; Pred. No. 0; Abtches 312; Conservative 0; Mismatches 0; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Overy Match 29.94: Score 485; DB 1: Louath 65:
Best Local Stmllarity 95.84; Pred No. 0;
Matches 91: Conservative 0; Mismiches 4; Indole 0: Capes
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APPLICANT: Lee James Beeley
APPLICANT: Lee James Beeley
APPLICANT: Robert James
TITLE OF INVENTION: NOVEL COMPOUNDS
FILE REPRENCE: GH-30002
CURRENT APPLICATION UNBERS: USG/09/142.563A
CURRENT FILM: DATE: 1999-06.30
EARLIER APPLICATION UNBERS: 9704551.2
EARLIER FILMS DATE: 1999-03-05
EARLIER FILMS DATE: 1999-03-18
EARLIER FILMS DATE: 1997-03-18
EARLIER FILMS DATE: 1997-03-18
EARLIER PILMS DATE: 1997-07-16
NUMBER OF SEQ ID MOS: 6
STOTHARE, PRASE 1997-07-16
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1. Sequence 6. Application US/09142565A
FALEN: No. 6187560
FORERAL INFORMATION:
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[1111111111]
301 LMKVOMLRESPF 312
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Sequence 6, Appli
Sequence 4, Appli
                                                                                                                                                                        September 29, 2003, 08:56:49; Search time 0.001 Seconds (without oil oilgment) 150:072 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. is the number of results predicted by chance to have a corre greater than or equal to the score of the result being printed, and is derived by analysis of the cotal score distribution.
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1620
1 WGLKPSDVPPTHAVKFLGA......TYEQLKRALMKVQMLRESPF 312
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1 1620 100.0 312 1 US-09-142-565-2 Sequence 2,
2 465 29.9 95 1 US-09-142-565-6 Sequence 6,
3 371 22.9 74 1 US-09-142-565-6 Sequence 6,
    GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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Sequence 2. Application US/09142565A
Patent No. 6197560
GENREAL NFORMATION
APPLICANT: Relly Patine
APPLICANT: Relly Patine
APPLICANT: Relly Patine
APPLICANT: Robert James
TITLE OF INVERTION: NOVEL COHOOUNDS
FILE REPERFORCE: GH-30002
CURRENT PILION DATE: 1999-06-00
EARLIER RILING DATE: 1999-06-00
EARLIER FILING DATE: 1999-01-01
EARLIER FILING DATE: 1997-01-01
EARLIER FILING DATE: 1997-01-16
EARLIER FILING DATE: 1997-01-16
EARLIER FILING DATE: 1997-01-16
SOOTHWARE: PRECEDO FOR MINGORE VERSION 3:0
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                                                                                                                       OM protein . protein search, using sw model
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Maximum Match 100%
Listing first 3 summaries
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Gapop 10.0, Gapext 0.5
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Maximum DB seq length: 200000000
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; ORGANISM: HOMO SAPIEN
US-09-142-565-2
                                                                                                                                                                                                                                                                                               Title:
Perfect score: ]
Sequence: ]
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JS-09-142-565-2
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1034 GCOACAGTGGTGGCCTCCCCCGGTGGACGTGCTGAACGCGCGTCTA 1093

[10] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [1] | [
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

18.94: Score 441.8: DB 1: Length 847:
Best Local Similarity 88.34 pred; No. O:
Bast Cocal Scimilarity 44: Wismatches 11: Indels 6: Gaps
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APPLICANT: Relly Patie
APPLICANT: Robert, James
APPLICANT: Robert, James
FILE & FERRING: ANNOREL COMPOUNDS
FILE BERRENCE: (1.000/141.565A
CURRENT FILING PARE: 1099-0.7442.565A
CURRENT FILING PARE: 1099-0.7445.12
EARLIER FILING PARE: 1999-0.7445.12
EARLIER FILING PARE: 1999-0.74551.2
EARLIER FILING PARE: 1997-0.716
EARLIER FILING PARE: 1997-0.716
EARLIER FILING PARE: 1997-0.716
SOFTWARE OF SEQ ID NOS: 6
SOFTWARE PARE: PARESCO FOR WINDOWS VORSION 3.0
SOFTWARE PARESCO FOR WINDOWS VORSION 3.0
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05-09-142-565-3
Sequence 3. Application US/09142565A
Patent No. 6457560
GENERAL INFORMATION:
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Sequence 5, Appli
Sequence 3, Appli
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No. Score Match Length DB ID
1 441.8 18.9 847 1 US-09-142-555-5
2 199.4 8.5 222 1 US-09-142-565-3
                                                                                           GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-142-56-5

Sequence 5, Application US/09142565A

Patent No. 61576-5

EARLEANT No. 6178-61

APPLICANT Lee James Beeley

APPLICANT Lee James Beeley

APPLICANT Lee James Beeley

APPLICANT ROBET James

TITLE 0F INVESTION: WOVEL COMPOUNDS

FILE REPRENCE: GH-30002

CURRENT APPLICATION UNDER: US/09/142,565A

CURRENT PILING DATE: 1939-06-30

EARLER PILING DATE: 1939-03-18

EARLER PAPLICATION UNDER: 1937-03-18

EARLER PILING DATE: 1937-03-18

EARLER PATE: 1937-03-18

EARLE
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                                                                                                                                                                                                                                                                                                        OM nucleic - nucleic search, using sw model
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Maximum Match 100%
Listing first 2 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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TYPE: DNA ORGANISM: HOMO SAPIEN US-09-142-565-3

TYPE: DNA
ORGANISM: HOMO SAPIEN
FEATURE:
NAME/KEY: unsure

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Search completed: September 29, 2003, 08:53:52 Job time : 2 secs

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Query Match 22.99; Score 371; DB 1; Length 74;
Best Local Similarity 93.21; Pred No. 0; Indels 0: Gaps 0;
Matches 69; Conservative 0; Mismatches 5: Indels 0: Gaps 0;
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Sequence 4. Application US/09142565A
Patent No. 618/7560
GENERAL INFORMATION:
APPLICANT: Hea James Beeley
APPLICANT: Heal James
APLICANT: Heal James
APLICANT: ROLE LANGER:
APLICANT: ROLE 1999-06-30
CURRENT APLICATION NOWER: US/09/141,565A
CURRENT AFLIKE DATE:
BARLIER APLICATION NOWER: US/05/14.7
EARLIER PILING DATE:
BARLIER FILING DATE:
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